

SEQUENCE LISTING

SEQ ID NO: 1

LENGTH: 377

TYPE: amino acid

5

TOPOLOGY: linear

MOLECULE TYPE: peptide

SEQUENCE DESCRIPTION:

	Met	Lys	Phe	Pro	Gly	Pro	Leu	Glu	Asn	Gln	Arg	Leu	Ser	Phe	Leu	
						5				10					15	
10	Leu	Glu	Lys	Ala	Ile	Thr	Arg	Glu	Ala	Gln	Met	Trp	Lys	Val	Asn	
						20				25					30	
	Val	Arg	Lys	Met	Pro	Ser	Asn	Gln	Asn	Val	Ser	Pro	Ser	Gln	Arg	
						35				40					45	
	Asp	Glu	Val	Ile	Gln	Trp	Leu	Ala	Lys	Leu	Lys	Tyr	Gln	Phe	Asn	
15						50				55					60	
	Leu	Tyr	Pro	Glu	Thr	Phe	Ala	Leu	Ala	Ser	Ser	Leu	Leu	Asp	Arg	
						65				70					75	
	Phe	Leu	Ala	Thr	Val	Lys	Ala	His	Pro	Lys	Tyr	Leu	Ser	Cys	Ile	
						80				85					90	
20	Ala	Ile	Ser	Cys	Phe	Phe	Leu	Ala	Ala	Lys	Thr	Val	Glu	Glu	Asp	
						95				100					105	
	Glu	Arg	Ile	Pro	Val	Leu	Lys	Val	Leu	Ala	Arg	Asp	Ser	Phe	Cys	
						110				115					120	
	Gly	Cys	Ser	Ser	Ser	Glu	Ile	Leu	Arg	Met	Glu	Arg	Ile	Ile	Leu	
25						125				130					135	
	Asp	Lys	Leu	Asn	Trp	Asp	Leu	His	Thr	Ala	Thr	Pro	Leu	Asp	Phe	

	140	145	150
	Leu His Ile Phe His Ala Ile Ala Val Ser Thr Arg Pro Gln Leu		
	155	160	165
	Leu Phe Ser Leu Pro Lys Leu Ser Pro Ser Gln His Leu Ala Val		
5	170	175	180
	Leu Thr Lys Gln Leu Leu His Cys Met Ala Cys Asn Gln Leu Leu		
	185	190	195
	Gln Phe Arg Gly Ser Met Leu Ala Leu Ala Met Val Ser Leu Glu		
	200	205	210
10	Met Glu Lys Leu Ile Pro Asp Trp Leu Ser Leu Thr Ile Glu Leu		
	215	220	225
	Leu Gln Lys Ala Gln Met Asp Ser Ser Gln Leu Ile His Cys Arg		
	230	235	240
	Glu Leu Val Ala His His Leu Ser Thr Leu Gln Ser Ser Leu Pro		
15	245	250	255
	Leu Asn Ser Val Tyr Val Tyr Arg Pro Leu Lys His Thr Leu Val		
	260	265	270
	Thr Cys Asp Lys Gly Val Phe Arg Leu His Pro Ser Ser Val Pro		
	275	280	285
20	Gly Pro Asp Phe Ser Lys Asp Asn Ser Lys Pro Glu Val Pro Val		
	290	295	300
	Arg Gly Thr Ala Ala Phe Tyr His His Leu Pro Ala Ala Ser Gly		
	305	310	315
	Cys Lys Gln Thr Ser Thr Lys Arg Lys Val Glu Glu Met Glu Val		
25	320	325	330
	Asp Asp Phe Tyr Asp Gly Ile Lys Arg Leu Tyr Asn Glu Asp Asn		

335 340 345
 Val Ser Glu Asn Val Gly Ser Val Cys Gly Thr Asp Leu Ser Arg
 350 355 360
 Gln Glu Gly His Ala Ser Pro Cys Pro Pro Leu Gln Pro Val Ser
 5 365 370 375
 Val Met

SEQ ID NO: 2

LENGTH : 1134

10 TYPE : nucleic acid

STRANDEDNESS : double

TOPOLOGY : linear

MOLECULE TYPE : DNA

SEQUENCE DESCRIPTION:

15 ATG AAG TTT CCA GGG CCT TTG GAA AAC CAG AGA TTG TCT TTC CTG 45
 TTG GAA AAG GCA ATC ACT AGG GAA GCA CAG ATG TGG AAA GTG AAT 90
 GTG CGG AAA ATG CCT TCA AAT CAG AAT GTT TCT CCA TCC CAG AGA 135
 GAT GAA GTA ATT CAA TGG CTG GCC AAA CTC AAG TAC CAA TTC AAC 180
 CTT TAC CCA GAA ACA TTT GCT CTG GCT AGC AGT CTT TTG GAT AGG 225
 20 TTT TTA GCT ACC GTA AAG GCT CAT CCA AAA TAC TTG AGT TGT ATT 270
 GCA ATC AGC TGT TTT TTC CTA GCT GCC AAG ACT GTT GAG GAA GAT 315
 GAG AGA ATT CCA GTA CTA AAG GTA TTG GCA AGA GAC AGT TTC TGT 360
 GGA TGT TCC TCA TCT GAA ATT TTG AGA ATG GAG AGA ATT ATT CTG 405
 GAT AAG TTG AAT TGG GAT CTT CAC ACA GCC ACA CCA TTG GAT TTT 450
 25 CTT CAT ATT TTC CAT GCC ATT GCA GTG TCA ACT AGG CCT CAG TTA 495
 CTT TTC AGT TTG CCC AAA TTG AGC CCA TCT CAA CAT TTG GCA GTC 540

CTT ACC AAG CAA CTA CTT CAC TGT ATG GCC TGC AAC CAA CTT CTG 585
CAA TTC AGA GGA TCC ATG CTT GCT CTG GCC ATG GTT AGT CTG GAA 630
ATG GAG AAA CTC ATT CCT GAT TGG CTT TCT CTT ACA ATT GAA CTG 675
CTT CAG AAA GCA CAG ATG GAT AGC TCC CAG TTG ATC CAT TGT CGG 720
5 GAG CTT GTG GCA CAT CAC CTT TCT ACT CTG CAG TCT TCC CTG CCT 765
CTG AAT TCC GTT TAT GTC TAC CGT CCC CTC AAG CAC ACC CTG GTG 810
ACC TGT GAC AAA GGA GTG TTC AGA TTA CAT CCC TCC TCT GTC CCA 855
GGC CCA GAC TTC TCC AAG GAC AAC AGC AAG CCA GAA GTG CCA GTC 900
AGA GGT ACA GCA GCC TTT TAC CAT CAT CTC CCA GCT GCC AGT GGG 945
10 TGC AAG CAG ACC TCT ACT AAA CGC AAA GTA GAG GAA ATG GAA GTG 990
GAT GAC TTC TAT GAT GGA ATC AAA CGG CTC TAT AAT GAA GAT AAT 1035
GTC TCA GAA AAT GTG GGT TCT GTG TGT GGC ACT GAT TTA TCA AGA 1080
CAA GAG GGA CAT GCT TCC CCT TGT CCA CCT TTG CAG CCT GTT TCT 1125
GTC ATG TAG 1134

15 SEQ ID NO : 3

LENGTH : 33

TYPE : nucleic acid

STRANDEDNESS : double

TOPOLOGY : linear

20 MOLECULE TYPE : DNA

SEQUENCE LISTING

CGTTCCCGGG TATGAAGTTT CCAGGGCCTT TGG 33

SEQ ID NO : 4

LENGTH : 31

25 TYPE : nucleic acid

STRANDEDNESS : double

TOPOLOGY : linear

MOLECULE TYPE : DNA

SEQUENCE LISTING

ACGGCTCGAG CTACATGACA GAAACAGGCT G

31

5

SEQ ID NO : 5

LENGTH : 14

TYPE : amino acid

TOPOLOGY : linear

10

MOLECULE TYPE : peptide

SEQUENCE LISTING

Glu Asp Asn Val Ser Glu Asn Val Gly Ser Val Cys Gly Thr

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